Thu Dec 12 16:02:06 2002

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COCATION: (14)
OTHER INFORMATION: Xaa = Leu or Val
NAME/KEY: WOD_RES
LOCATION: (16)
OTHER INFORMATION: Xaa = Met or Leu
NAME/KEY: WOD_RES
LOCATION: (17)
OTHER INFORMATION: Xaa = Glu or Asp
NAME/KEY: WOD_RES
LOCATION: (18)
OTHER INFORMATION: Xaa = any amino acid, Xaa at positions 18 and 19
OTHER INFORMATION: may be present or absent
                      APPLICANT: Fischer, Robert L.
APPLICANT: Fischer, Robert L.
APPLICANT: Choi, Yeonhee
APPLICANT: Hannon, Mike
APPLICANT: Hannon, Mike
APPLICANT: Hannon, Mike
APPLICANT: Tatarinova, Jack Kishiro
APPLICANT: Tatarinova, Tatiana Valerievna
APPLICANT: The Regents of the University of California
TITLE OF INVENTION: Nucleic Acids That Control Plant Development
FILE REFERENCE: 023070-099910US
CURRENT FILING DATE: 2001-04-23
PRIOR APPLICATION NUMBER: US/09/840,743
PRIOR PLILNG DATE: 2000-04-21
NUMBER OF SEQ ID NOS: 119
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 71
LENGTH: 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Description of Artificial Sequence:DMT Domain A consensus sequence
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NAME/KEY: MOD_RES
LOCAPTON: ....
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OTHER INFORMATION: Xaa = any amino acid
MAME/KEY: MOD_RES
LOCATION: (4)
OTHER INFORMATION: Xaa = 11e or Leu
NAME/KEY: MOD_RES
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NAME/KEY: MOD_RES
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OTHER INFORMATION: Xaa = Asp or Pro
NAME/KEY: MOD_RES
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LOCATION: (7)
OTHER INFORMATION: Xaa = Glu or Val
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Sequence 71, Application US/09840743 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Artificial Sequence
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OTHER INFORMATION: C
OTHER INFORMATION: C
NAME/KEY: MOD_RES
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LOCATION: (21)
OTHER INFORMATION: X
NAME/KEY: MOD_RES
LOCATION: (22)
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OTHER INFORMATION:
NAME/KEY: MOD_RES
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NAME/KEY: MOD_RES
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ION: (28)
INFORMATION: Xaa = any amino acid
                                                                          NAME/KEY: MOD_RES
LOCATION: (32). (33)
OTHER INFORMATION: Xaa = any amino acid
NAME/KEY: MOD_RES
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OTHER INFORMATION: Xaa = any amino acid
NAME/KEY: MOD_RES
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ON: (27)
INFORMATION: Xaa = Trp or Leu
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NAME/KEY: MOD_RES
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NAME/KEY: MOD_RES
LOCATION: (87)
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NAME/KEY: MOD_RES
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NAME/KEY: MOD_RES
LOCATION: (56)
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LOCATION: (55)
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LOCATION: (74)
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LOCATION: (12)..(17)

OTHER INFORMATION: Xaa — any amino acid, Xaa at positions 15-17 may be OTHER INFORMATION: present or absent
NAME/KEY: MOD_RES
LOCATION: (19)
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APPLICANT: Tatarinova, Tatlana Valerievna
APPLICANT: The Regents of the University of California
TITLE OF INVENTION: Nucleic Acids That Control Plant Development;
FILE REFERENCE: 023070-099910US
CURRENT APPLICATION NUMBER: US/09/840,743
PRIOR APPLICATION NUMBER: US/09/840,743
PRIOR APPLICATION NUMBER: US/09/840,743
PRIOR RILING DATE: 2000-04-21
NUMBER OF SEQ ID NOS: 119
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 72
LENGTH: 230
TYPE: PRT
CRANISM: Artificial Sequence
FEATURE:
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NAME/KEY: MOD_RES
                                                                                                                                                  OTHER INFORMATION: Xaa - any amino acid
NAME/KEY: MOD_RES
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NAME/KEY: MOD_RES
LOCATION: (21)
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OTHER INFORMATION: Xaa = any amino acid
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INFORMATION: Xaa = any amino acid
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INFORMATION: Xaa = any amino acid
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OTHER INFORMATION: Xaa = any amino acid
OTHER INFORMATION: Xaa = Ala or Ser
NAME/KEY: MOD_RES
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NAME/KEY: MOD_RES
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APPLICANT: Fischer, Robert L.
APPLICANT: Choi, Yeonhee
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OTHER INFORMATION: Xaa = any amino acid
NAME/KEY: MOD_RES
LOCATION: Xaa = any amino acid, Xaa at positions 53-62 may be
OTHER INFORMATION: present or absent
NAME/KEY: MOD_RES
LOCATION: (65)..(67)
OTHER INFORMATION: Xaa = any amino acid
NAME/KEY: MOD_RES
LOCATION: (65)..(77)
OTHER INFORMATION: Xaa = any amino acid
NAME/KEY: MOD_RES
LOCATION: (69)..(70)
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COCATION: (71)
OTHER INFORMATION: Xaa = His or Asn
NAME/KEY: MOD_RES
COCATION: (73)
OTHER INFORMATION: Xaa = any amino acid
NAME/KEY: MOD_RES
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OTHER INFORMATION: Xaa = any amino acid
NAME/KEY: MOD_RES
                                          LOCATION: (44)..(45)
OTHER INFORMATION: Xaa = any amino acid
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NAME/KEY: MOD_RES
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OTHER INFORMATION: Xaa = Asn or Phe NAME/KEY: MOD_RES
                                                                                NAME/KEY: MOD_RES
LOCATION: (47)
OTHER INFORMATION: Xaa = Ala or Ser
NAME/KEY: MOD_RES
LOCATION: (48)
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OTHER INFORMATION: Xaa = His or Lys
NAME/KEY: MOD_RES
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NAME/KEY: MOD_RES
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NAME/KEY: MOD_RES
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OTHER INFORMATION: X NAME/KEY: MOD_RES LOCATION: (109)
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OTHER INFORMATION: NAME/KEY: MOD_RES
LOCATION: (89)
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NAME/KEY: MOD_RES
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NAME/KEY: MOD_RES
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OTHER INFORMATION:
NAME/KEY: MOD_RES
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LOCATION: (84)
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LOCATION: (95)
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LOCATION: (212)..(221)
OTHER INFORMATION: Xaa = any amino acid, Xaa at positions 212-221 may
OTHER INFORMATION: be present or absent
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NAME/KEY: MOD_RES
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OTHER INFORMATION: Xaa - any amino acid
NAME/KEY: MOD_RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    - any amino acid
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US-09-840-743-72
                                         - Arg or Lys
                                                                                           His or Tyr
                                                                                                                                                - Phe or Tyr
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OTHER INFORMATION: Xaa - Ala or Ser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION: Xaa - Arg or Lys
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                                                                                                                                                                                                      = Ala or Ser
                                                                                                                                                                                                                                                                                           LOCATION: (211)
OTHER INFORMATION: Xaa = Ala or Val
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      - Ala or
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NAME/KEY: MOD_RES
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INFORMATION: Xaa
                                       INFORMATION: Xaa
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OTHER INFORMATION:
NAME/KEY: MOD_RES
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LOCATION: (207)
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OTHER INFO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LOCATION: (183)..(184)
OTHER INFORMATION: Xaa = any amino acid, Xaa at positions 183 and 184
OTHER INFORMATION: may be present of absent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LOCATION: (150)
OTHER INFORMATION: Xaa = any amino acid
NAME/KEY: MOD_RES
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OTHER INFORMATION: Xaa - any amino acid
NAME/KEY: MOD_RES
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                    - any amino acid
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LOCATION: (152)
OTHER INFORMATION: Xaa = Glu or Asp
NAME/KEY: MOD_RES
'``ATTION: (153)
'``ATTION: (153)
                                                    LOCATION: (125)
OTHER INFORMATION: Xaa - Ala or Cys (-
                                                                                                                                                                                                                                                                                                                                                                                                                                 LOCATION: (144)
OTHER INFORMATION: Xaa = Leu or Gln
NAME/KEY: MOD_RES
OUGHER INFORMATION: Xaa = Glu or Phe
NAME/KEY: MOD_RES
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OTHER INFORMATION: Xaa = 11e or Val
NAME/KEY: MOD_RES
OTHER INFORMATION: (166)
OTHER INFORMATION: Xaa = Asp or Pro
NAME/KEY: MOD_RES
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OTHER INFORMATION: Xaa - Gln or Glu
NAME/KEY: MOD_RES
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OTHER INFORMATION: Xaa = Leu or Val
NAME/KEY: MOD_RES
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NAME/KEY: MOD_RES
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NAME/KEY: MOD_RES
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OTHER INFORMATION: Xaa - Leu or Met
                                                                                                                                                                                  - Trp or Leu
                                                                                                                                                                                                                                                                                                                                                                                                   or Met
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INFORMATION: Xaa = Leu or Met
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                                                                                                                             - Met or
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LOCATION: (142)
OTHER INFORMATION: Xaa - Leu
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LOCATION: (130)
OTHER INFORMATION: Xaa
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OTHER INFORMATION: Xaa
NAME/KEY: MOD_RES
LOCATION: (176)
LOCATION: (112)..(113)
OTHER INFORMATION: Xaa
                                                                                                        LOCATION: (128)
OTHER INFORMATION: Xaa
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OTHER INFORMATION: Xaa
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OTHER INFORMATION: Xaa
                                                                                         MOD_RES
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LOCATION: (189)...
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LOCATION: (183)..
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LOCATION:
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